

## Untangling the reticulate history of species complexes and horticultural breeds in *Abelia* (Caprifoliaceae)

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1 original article

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5 **Untangling the reticulate history of species complexes and**  
6 **horticultural breeds in *Abelia* (Caprifoliaceae)**

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## ABSTRACT

**• Background and Aims** The genetic and morphological consequences of natural selection and selective breeding are explored in the genus *Abelia*. The genus consists of ornamental shrubs endemic to China, which have been bred to create attractive and diverse cultivars.

**• Methods** We use DNA fingerprinting (AFLP) and DNA sequence data to investigate the genetic diversity among 46 accessions of *Abelia* (22 natural taxa and 24 horticultural breeds). In the cultivated varieties these data are used to explore taxon boundaries, hybridisation and backcrossing. The dataset is also used to investigate morphological variation within natural species complexes and subsequently to inform a taxonomic treatment.

**• Key Results** *Abelia* comprises five species: *A. forrestii*, *A. schumannii*, *A. macrotera*, *A. uniflora* and *A. chinensis* and has a total of 11 varieties. *Abelia uniflora* and *A. macrotera* do not occur in sympatry and are disjunctly distributed to the east and west of the *A. chinensis* distribution range. *Abelia chinensis* is widespread in eastern China and creates hybrids and introgressive taxa, including *A. uniflora*, along the contact zones with the previous taxa. *Abelia* ‘Maurice Foster’ is a horticultural variety collected from wild stocks in Sichuan (China). Bayesian clustering methods (inferred in STRUCTURE based on AFLP data) indicate admixture between *A. macrotera* and *A. schumannii* in this variety. We can infer that hybridization probably occurred in the wild where these progenitor taxa co-occur and naturally form hybrids. AFLP results also reveal that a few diagnostic morphological characters such as sepal number or inflorescence structure were transferred between natural species are mirrored by horticultural crosses such as in *Abelia* ‘Saxon Gold’ and *A. forrestii*.

**• Conclusions** Studying both natural and cultivated species from the same group has enabled us to understand both differentiation mechanisms and how to improve cultivated plants in the future by studying which morphological characters are transferred between species and which taxa may already have arisen through hybridisation.

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**Keywords:** *Abelia*; AFLP; hybridization and introgression; cultivars; nomenclature; China.

**Short Title:** *Abelia* hybridization and introgression

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## INTRODUCTION

Described by Robert Brown in 1818 ([Brown, 1818](#)) the genus *Abelia* is known for its widely cultivated hybrid, *A. x grandiflora*, commonly cultivated as a trimmed hedge around the world. Although its breeding history is documented, exactly how it relates to the wild taxa occurring in China remains unclear. The hybrid is said to have been obtained in Italy by MM. Rovelli Brothers Nurseries in Pallanza (Lake Maggiore), a cross between *Abelia chinensis* and *A. uniflora* ([André, 1886](#)). *Abelia* is a genus endemic to China with between three and 15 taxa, depending upon which circumscription is followed. It is regarded as a species complex with difficult taxonomic treatment. Most specimens of *Abelia* examined are diploid (Kim, 1998) and probably self-incompatible (Scobie and Wilcock, 2009), they do not vegetatively reproduce but are propagated clonally by cuttings in cultivation. Viable seed production is low and achenes are wind dispersed, though dispersal distances have not been studied. The current study aims to compare and contrast the genetic signature of selective breeding and genetic differentiation in the genus *Abelia*. To improve the feasibility of this study we first needed to have a clear idea of the morphological variation, distribution and nomenclature of the naturally occurring taxa of *Abelia* in China as well as those in cultivation. We have devised a new nomenclature and studied in detail the morphological and genetic variation within *Abelia* and how selective breeding and hybridisation could give answers to the species concept.

### *Mechanisms of differentiation*

Polyploidization and hybridization are among the most important forces in the evolution of higher plants and at least 25% of the plant species are involved in hybridisation and introgression with other species ([Mallet, 2005, 2007](#)). Hybrids combine different genotypes and generate phenotypic traits that are often intermediate between their parents and

1 that in turn is often used as evidence for hybridization in morphological analyses. If  
2 backcrossing with one or both the parental taxa occurs repeatedly, i.e. introgression occurs,  
3 the parental taxa successively incorporate parts of the genome from the other taxa involved in  
4 hybridization and further increase the number of morphological transitions between the  
5 species as demonstrated in other studies ([Rieseberg, 1997](#); [Hardig et al., 2000](#)). Uncovering  
6 hybridization and introgression is important to reveal the origin of species and to obtain  
7 insights into the processes behind the intra- and interspecific variability.

8 Interspecies hybridization is a method that is often used in horticultural breeding  
9 programmes, wild relatives are used as donors of desirable traits for the cultivated plants (for  
10 example disease resistance). Provenance and sometimes identity of the parents are rarely  
11 documented, this problem is more acute in groups of species or species complexes where taxa  
12 are morphologically similar and difficult to distinguish: ‘Some of the confusion in the naming  
13 of cultivated material appears to derive from a failure to recognise the great variability of  
14 some wild taxa’ ([Barnes, 2001](#)). Breeding programmes are nevertheless a model to study  
15 hybridisation and its consequences because artificial hybrids unlike wild populations are not  
16 involved in uncontrolled introgression or selection which would blur limits between taxa.

17 Cultivars and hybrids could therefore be a model to trace back the crossing history of wild  
18 taxa and the consequences on their phenotypic traits. Incomplete lineage sorting is  
19 nevertheless also to be taken into account.

20 One approach for identifying hybrids uses incongruences between phylogenies based on  
21 maternally (cpDNA) and nuclear genes inherited markers ([Cronn et al., 2003](#); [Clarkson et al.,](#)  
22 [2010](#)). A premise is that the phylogenetic resolution of the studied markers is sufficient to  
23 distinguish taxa suspected to be involved in hybrid formation. This is often not met in plastid  
24 DNA regions which generally have low substitution rates ([Wolfe et al., 1987](#)). In recent  
25 years, a number of molecular markers, e.g., random amplified polymorphic DNA (RAPD;

1 (Williams *et al.*, 1990)), amplified fragment length polymorphism (AFLP; (Vos *et al.*, 1995))  
2 simple sequence repeats (SSR; (Zietkiewicz *et al.*, 1994)) and inter-simple sequence repeats  
3 (ISSR) have been widely used to detect genetic diversity in plants (Nybon and Bartish, 2000).  
4 Various other ornamental plants have been used for DNA marker based diversity studies  
5 using AFLP including *Nelumbo* (Hu *et al.*, 2012), *Rosa* (Koopman *et al.*, 2008), yellow  
6 Camellia (Tang *et al.*, 2006). AFLP was used in studies of two related *Abelia* genera,  
7 *Diabelia* in Zhejiang (Zhou *et al.*, 2004) and *Dipelta* in Gansu (Liu *et al.*, 2013).

## 8 MATERIALS AND METHODS

### 9 *Sampling*

10 A total of 55 samples were collected mostly from the trial field experiment at RHS  
11 Wisley (UK) spanning the morphological and geographical variability within targeted taxa  
12 (Appendix 1).

13

### 14 *DNA extraction*

15 Genomic DNA was extracted using a modified cetyltrimethylammonium bromide  
16 (CTAB) method (Doyle and Doyle, 1987) using 0.2 g of silica dried leaf tissue. A standard  
17 protocol was used to obtain purified DNA (Mikulášková *et al.*, 2012).

18

### 19 *ITS and cpDNA sequences*

20 Published and unpublished sequences were gathered from two previous studies by  
21 Landrein *et al.* (2012) and Hua Feng *et al.* (2014). See publications for detailed methods and  
22 Table 1 for gene Bank numbers. *Vesalea floribunda* was selected as an outgroup based on  
23 previous data. Phylogenetic analyses were performed using PAUP\* v4b10 (Swofford, 2002)  
24 for maximum parsimony (MP) analyses, and MrBayes 3.2.2 (Ronquist and Huelsenbeck,

1 2003) for Bayesian inference (BI). The MP analyses used heuristic searches with 1,000  
2 random addition sequence replicates, tree bisection reconnection (TBR) branch swapping,  
3 and MULTREES on. All character states were treated as unordered and equally weighted  
4 with gaps treated as missing data. To evaluate the relative robustness of clades in the MP  
5 trees, a bootstrap analysis (Felsenstein, 1985) was performed with 1,000 replicates using the  
6 same options as above except that a maximum of 100 trees were saved per replicate.  
7 MrModeltest 2.3 (Nylander, 2004) was run for each of the data sets to determine the most  
8 appropriate substitution models using the Akaike criterion (Posada and Buckley, 2004) (see  
9 the final row of Table 1). To estimate support for each node, 1 000 bootstrap replicates were  
10 performed with automatic termination at 10,000 generations, All final runs were performed  
11 on the CIPRS Science Gateway (<http://www.phylo.org/portal2/>) (Miller *et al.*, 2010). A  
12 partitioned Bayesian analysis of the plastid dataset was also implemented by applying the  
13 previously determined models to each data partition (Brown and Lemmon, 2007). ITS was  
14 found to be incongruent with the plastid analysis and the two were not combined as shown in  
15 Landrein *et al.* (2012) and Wang *et al.* (2015). For BI 40 million generations were run with  
16 four chains, each starting with a random tree. Trees were sampled every 1,000 generations.  
17 Posterior probabilities (PP) were calculated from the majority consensus of all the sampled  
18 trees. When the standard deviation of the split frequencies (SDSF) permanently fell below  
19 0.01, 10% of the trees sampled during the burn-in phase were discarded. All final runs were  
20 performed on the CIPRS Science Gateway (<http://www.phylo.org/portal2/>) (Miller *et al.*,  
21 2010).

22

### 23 *AFLP amplification and scoring*

24 Reactions were performed following the protocol of Arrigo *et al.* (2010). After  
25 performing a primer trial with 24 primers, three selective primer pairs were chosen (*EcoR1*-



1 ACG/Mse1-CTA; *Eco*R1-AAG/Mse1-CAC and *Eco*R1-ACC/Mse1-CTG), with JOE and  
2 NED-labelled *Eco*R1 primers. PCR products were mixed with a Rox size standard ladder and  
3 analysed with an ABI 3730XL capillary sequencer. In order to detect and calculate the size of  
4 AFLP bands, raw electropherograms were analysed using Peakscanner (ABI) with default  
5 parameters except a light peak smoothing. A binary matrix of AFLP band presence (1) and  
6 absence (0) was built using the automated scoring RawGeno package (R CRAN; ([Arrigo et](#)  
7 [al., 2009](#))) with the following parameters: scoring range, 50-400 bp; minimum intensity, 100  
8 rfu; minimum bin width, 1 bp; maximum bin width, 1.5 bp. Closely sized bins were  
9 eliminated.

10 Individuals were randomly distributed in 96-well plates in order to produce a reliable AFLP  
11 dataset. 53 samples (representing 22% of the final dataset) were randomly chosen from each  
12 plate and replicated to calculate the error rate ([Bonin et al., 2004](#)). Bands that were clearly  
13 not reproducible were discarded from further analysis. Population structure was analysed  
14 with STRUCTURE v.2.3.2.1 ([Pritchard et al., 2000](#)) using an admixture model. Bayesian  
15 estimates of genetic clustering probabilistically assigns individuals to populations defined by  
16 allele frequencies at multiple loci. (Pritchard & al 2000). Parameters selected were: 5000  
17 burn-in repetitions and 50000 MCMC simulations at four iterations.

18 Our *Abelia uniflora* extraction was unsuccessful with AFLP protocols, as this requires higher  
19 molecular weight DNA but this species was included in the DNA sequencing analyses.

20 To assess the best *K* for the results from STRUCTURE were analysed in CLUMPAK (a  
21 program for identifying clustering modes and packaging population structure inferences  
22 across *K* by Kopelman, Naama M; Mayzel, Jonathan; Jakobsson, Mattias; Rosenberg, Noah  
23 A; Mayrose) according to the Evanno method ([Evanno et al., 2005](#)).

24

25

## RESULTS

### *CpDNA-based phylogeny*

For 24 taxa, the number of characters was 2232, 321 (14.4%) of which were variable and 168 (7.5%) of which were potentially parsimony informative. The number of characters contributed by each individual region was 604 from *rbcL*, 849 from *trnL-F* and 779 from *matK*. Details for each region are provided in Table 1.

*Abelia x grandiflora* and *Abelia chinensis* form a clade (BP 54/ PP 0.96, Fig. 1) and a polytomy with *A. macrotera* var. *deutziaefolia*. Both varieties of *Abelia forrestii* form a clade (BP 99, PP 1, Fig. 1) but there is no resolution for the rest of the taxa. *A. x grandiflora* from Zhejiang is sister to *Abelia x grandiflora* originating in Europe (BP 78/ PP 1 Fig. 1).

### *ITS*

For 18 taxa, the number of characters was 637, 65 (10.2%) of which were variable and 40 (6.2%) of which were potentially parsimony informative. Details are provided in Table 2. The monophyly of the genus *Abelia* is weakly supported (PP 0.57, Fig. 2) and *Abelia uniflora* forms a clade with *A. x grandiflora* (BP 91/ PP 1, Fig. 2). *Abelia schumannii* and *A. macrotera* form a separate clade (BP 78/ PP 1).

### *AFLP*

Replicate samples indicated high reproducibility of the AFLP data, with the error rate being 0.0%. Altogether, 488 AFLP markers were scored; 300 (61%) of these markers were polymorphic. Details for each region are provided in Table 2.

A Neighbour Joining Tree (Fig. 3) and PCoA (Fig. 4) revealed four clusters that correspond to individuals of

1. *Abelia chinensis* and backcrosses
2. *Abelia macrotera*
3. *Abelia schumannii*, *A. macrotera* x *schumannii* ‘Maurice Foster’, *A.* ‘Saxon Gold’  
and *A.* ‘Edward Goucher’
4. *Abelia* x *grandiflora* and *A. forrestii*

*Abelia forrestii* is sister to *A. macrotera* and *A. schumannii* but clusters with *A. x grandiflora* in the PCoA.

Within *A. x grandiflora* genetic variability was low and three groups can be identified:

1. *Abelia* x *grandiflora* ‘Sherwood’
2. *Abelia* x *grandiflora* ‘Francis Mason’
3. *Abelia* x *grandiflora* ‘Little Richard’

Dwarf, variegated plants can be found in all three groups. ‘Sherwood’-like flowers with basal spurs and dark leathery leaves can also be found in two groups ‘Sherwood’ and ‘Little Richard’.

The results generated by STRUCTURE give some indication on the origin of *A. x grandiflora* and admixture of alleles. The analysis provided strongest support for  $K = 2$  when considering  $\Delta K$  and  $K = 4$  when considering  $\text{LnP}(\text{Pr data})$ . We identified a substructure at  $K=5$  in which populations appeared homogeneous in their admixture composition. Higher values of  $K$  yielded noise that appeared as ancestry shared by very few individuals within the same populations. Cultivated taxa with fully traceable breeding histories such as *A.* ‘Edward Goucher’ (*A. x grandiflora* x *schumannii*) were also used to verify the best  $K$ .

The five populations identified are (Fig. 5):

1. *Abelia macrotera*
2. *Abelia schumannii*
3. *Abelia forrestii*

1        4. *Abelia chinensis*

2        5. *Abelia x grandiflora*

3        Admixture was identified in *A. macrotera x schumannii* ‘Maurice Foster’, wild collected in  
4        Sichuan, at a contact zone between the two taxa. Admixture is also shown with *A.* ‘Edward  
5        Goucher’ as known from the breeding records, as well as in *A.* ‘Saxon Gold’ with *A.*  
6        *schumannii*, which had not been recorded previously. Backcrosses with *A. chinensis*  
7        (‘Canyon Creek’; ‘Pleasant surprise’; ‘Rose Creek’) also show admixture with *A. chinensis* as  
8        informed by the breeding records and the results shown here. No admixture was identified for  
9        *A. x grandiflora* suggesting one of the parents was not present in the sampling or the parents  
10       are themselves also the result of hybridisation and introgression in the past (Baack and  
11       [Rieseberg, 2007](#)).  
12

13       *Abelia morphological study and nomenclature*

14       The AFLP analysis (detailed previously) identified 5 natural taxa in *Abelia*. However,  
15       these taxa do not correspond with the current taxonomy and therefore there is a need to re-  
16       examine morphological characters and make some refinements to the classification for the  
17       group. The current taxonomy for the genus was last revised by Rehder (Rehder 1911) and  
18       was useful in assessing the morphological diversity within the group. Some characters can be  
19       misleading especially in many cryptic taxa which are sometimes morphologically distinct, but  
20       are difficult to assess because their genetic histories are unknown.

21       Here are presented the results of an analysis of herbarium specimens from 16 herbaria (A,  
22       BM, CAS, CDBI, E, GXMI, HENU, HIB, IBK, IBSC, K, KUN, LBG, P, PE, W: acronyms  
23       according to Holmgren *et al.* (1990)), published literature sources, including all of the  
24       Chinese regional Floras and field observations. A full revision of the genus will be published  
25       separately (Landrein in prep).

1 *Abelia* was first monographed by Graebner (1901), 10 species were recognised, three more  
2 species were added by Rehder (1911) and this was reduced to only five species in *Flora*  
3 *Sinica* (Hsu *et al.*, 1988). A total of 25 names have been published.

4 In this study we recognise five species and 11 varieties. Morphological variation between  
5 specimens (761 specimens studied) could mostly be attributed between only two character  
6 states such as bilabiate versus infundibuliform corollas; white versus pink corollas; scented  
7 versus non-scented corollas; orange markings versus no markings on corolla mouth; long  
8 exerted versus inserted stamens, five calyx lobes versus two calyx lobes, paired versus  
9 single flowers, four versus six episepals, loose versus compact inflorescences. Characters  
10 were nevertheless randomly distributed within species complexes such as *Abelia forrestii*  
11 with five sepals instead of two or *Abelia uniflora* with a variable number of sepals. A few  
12 apomorphic characters were also identified such as the inflated mouth corolla in *A.*  
13 *schumannii* (Table 3, Figs. 6 & 7).

14

#### 15 *Nomenclature of taxa*

16 see appendix 2

17

#### 18 *Key*

19 see appendix 3

20

#### 21 *Distribution*

22 *Abelia* is widely distributed across much of China. Its range is restricted in the north by the  
23 Qin Ling Range, in the west by the deep valleys of the Qinghai-Tibetan plateau (QTP), in the  
24 east by the East China Sea reaching the Ryukyu Islands and to the South reaching the  
25 Mountains of Northern Vietnam. The species are distributed in a wide array of habitats

(roadside, woodlands, and scrublands) restricted to mountains, deep valleys and hills gradually diversifying from West to East. Most taxa are widespread with overlapping ranges, but some can be localised and with restricted ranges (Fig. 8).

- *Abelia forrestii* var. *forrestii* is restricted to one locality in the Nujiang valley.
- *Abelia forrestii* var. *gracilentia* is restricted to the first bend of the Yangtze River in the Leaping Tiger Gorge.
- *Abelia macrotera* var. *zabelioides* is confined to E'mei Shan but intermediate forms are more widespread.
- *Abelia macrotera* var. *parvifolia* is only recorded from around Yichang in Hubei.
- *Abelia macrotera* var. *myrtilloides* has a small range in northern Sichuan.
- *Abelia chinensis* var. *achersoniana* is only found in Hong Kong and may be a coastal ecotype.
- *Abelia chinensis* var. *lipoensis* can be found in Southern Yunnan, Guizhou, Guangxi and Guangdong.
- *Abelia chinensis* var. *hanceana* grows in Fujian: Xiamen and also Taiwan as well as the Ryukyu Islands.
- *Abelia uniflora* grows in Fujian, Jiangxi, Anhui and Zhejiang

#### *Cultivars*

The breeding of *Abelia* only started in 1950 more than 50 years after the discovery of the hybrid *Abelia* x *grandiflora*. The origin of *Abelia* x *grandiflora* has been documented by Ed. André (1886) 'Seen in the M.M. Rovelli Brothers Nurseries in Pallanza (Lake Maggiore); grown from a seedling a few years ago this shrub, that MM. Rovelli named (*A. rupestris grandiflora*) is more vigorous with more persistent leaves, ovate, serrate, glossy green and with large tubular flowers, fragrant, pinkish rose, blooming during all seasons, in full sun'.

1 Its morphology strongly suggests it is of hybrid origin between *A. chinensis* R.Br. (syn:  
2 *Abelia rupestris* Lindl.) and *Abelia uniflora*. The morphological characters are intermediate  
3 between the two parents. Flowers are axillary or in small cymes, single and with 4 episepals  
4 similar to *A. uniflora*. The flowers are white, fragrant, infundibuliform-campanulate and  
5 slightly bilabiate, the stamens are slightly exerted and therefore intermediate between the  
6 two parents. The leaves are more similar to *A. chinensis* (Table 4, Figs. 9 & 10).  
7 For several decades breeding was focused on obtaining yellow, variegated leaves as well as  
8 dwarf variants. Recent breeding has focused on new hybridisation including *A. schumannii*  
9 and *A. chinensis* var. *chinensis* and the creation of several backcrosses that are of interest in  
10 this study. Amongst them we can cite 'Saxon Gold', and 'Rose Creek'. The first backcross  
11 recorded nevertheless predates this recent activity and 'Edward Goucher', one of the most  
12 popular cultivars, was obtained in 1911. Michael Dirr has been an active participant in most  
13 breeding programs at the University of Georgia (Dirr, 2009).  
14 'Francis Mason' was obtained in 1950 in the Mason nurseries, New Zealand. It is by far the  
15 most commonly grown cultivar of *Abelia* x *grandiflora* and is distinguished by its yellow  
16 variegated leaves often reverting to all yellow or green. Another origin of *A. x grandiflora*  
17 cultivars is 'Sherwood' published in the Proceedings at the annual meeting of the American  
18 Association of Nurserymen, Florists and Seedsmen 1949:123, it has a more compact habit. A  
19 few variegated forms have also been obtained from 'Sherwood'. See Table 5 for list of  
20 hybrids and cultivars with their breeding history.

21

## 22 *Morphological characters*

23 Morphological variation between cultivars was paralleled with the variation in wild taxa and  
24 could mostly be attributed between two character states such as glossy versus non glossy  
25 leaves; bilabiate versus infundibuliform corollas; white versus pink corollas; scented versus

1 non scented corollas; long exerted versus inserted stamens, five calyx lobes versus two calyx  
2 lobes, paired versus single flowers, four versus six episepals, loose versus compact  
3 inflorescences. Characters were randomly distributed within hybrids and backcross such as  
4 *Abelia* ‘Saxon Gold’ with five sepals instead of two or *Abelia x grandiflora* with a variable  
5 number of sepals. A few apomorphic characters were also identified such as the variegated  
6 leaves or corolla spur in *Abelia* ‘Sherwood’ (Table 4, Figs. 9 & 10).

7

## 8 DISCUSSION

9 *Abelia* is a typical genus of the Sino-Japanese Floristic Region (SJFR) (Qiu *et al.*,  
10 2011). The QTP is the highest and largest plateau in the world and is regarded as a  
11 biodiversity hotspot where deep valleys have allowed for many species to occur  
12 sympatrically, however hybridisation and introgression are common problems for species  
13 delimitation (Xu *et al.*, 2010; Liu *et al.*, 2013). A clear West-East distribution pattern can be  
14 observed with *A. macrotera* var. *mairei* in the most western part (Yunnan, Hengduan  
15 Mountains). *Abelia schumannii* is restricted to the deep valleys of the QTP. *Abelia forrestii* is  
16 the most localised species and restricted along the Salween-Mekong divide (Nujiang River  
17 (Salween) for var. *forrestii* and the first bend of the Yangtze River for var. *gracilentia*). *Abelia*  
18 *chinensis* is distributed in a wide area in eastern China as well as more isolated populations in  
19 Hubei and Sichuan. *Abelia chinensis* diversifies at the contact of *Abelia macrotera* and *A.*  
20 *uniflora* with *A. chinensis* var. *lipoensis* in South Yunnan, Guangxi and Guizhou and *A.*  
21 *chinensis* var. *hanceana* in Xiamen, Taiwan and the Ryukyu Islands. Finally, a clear  
22 disjunction can be observed between *A. uniflora* in Anhui, Fujian, Jiangxi and Zhejiang with  
23 *Abelia macrotera* in Hubei and further west; this seems to be linked with the dominance of *A.*  
24 *chinensis* in Jiangxi, Hunan and Guangxi. A north-south distinction can be observed with *A.*



1 *macrotera* var. *deutziaefolia* occurring in Yunnan and Guangxi which may also been driven  
2 by the presence of *A. chinensis* in the same area (Fig. 8).

3 Possible phylogeographic events cited from the literature:

- 4 1. *Abelia macrotera* - *Abelia uniflora* west-east disjunction with allopatric  
5 incipient speciation (Qiu *et al.*, 2009a)
- 6 2. *Abelia schumannii* range expansion from SE plateau edge onto eastern QTP  
7 platform (Yang *et al.*, 2008)
- 8 3. *Abelia forrestii* glaciation survivor east and west of the Mekong-Salween  
9 divide (Li *et al.*, 2011)
- 10 4. *Abelia macrotera* var. *deutziaefolia* and *A. chinensis* var. *lipoensis* genetic  
11 differentiation among isolated populations from Guangxi, Guangdong, Hunan  
12 and Hainan; localized range expansion, possibly during interglacial periods  
13 (Tian *et al.*, 2010)
- 14 5. *Abelia chinensis* var. *hanceana* ancient allopatric-vicariant segregation across  
15 Japan vs. eastern China (Qiu *et al.*, 2009b)

16

#### 17 ***Abelia macrotera* – *Abelia uniflora*: Hybridization and introgression with *Abelia chinensis***

18 *Abelia uniflora* is the most enigmatic species in the genus *Abelia*, it was only briefly  
19 described (Wallich, 1829) and was the first taxon to be introduced into cultivation (as early as  
20 1845 by Robert Fortune). It was illustrated in the *Curtis Botanical Magazine* (Hooker, 1853)  
21 and is likely to have become extinct in cultivation after this date. *Abelia* x *grandiflora* is said  
22 to have been obtained in Italy by MM. Rovelli Brothers Nurseries in Pallanza (Lake  
23 Maggiore) as a cross between *Abelia chinensis* and *A. uniflora* (André, 1886).

1 The ITS sequence analysis (Fig. 2) shows that *Abelia uniflora* forms a clade with *A. x*  
2 *grandiflora* (BP 91/ PP 1, Fig. 2) but the position is unresolved alongside *A. macrotera* in the  
3 cpDNA analysis. This is an indication of the hybrid origin of cultivated *Abelia x grandiflora*  
4 involving *A. uniflora* as one progenitor but the other progenitor is not identified definitively.  
5 It may be *A. chinensis* as stated in the literature or *A. macrotera* as indicated (albeit weakly)  
6 in the phylogenetic analysis or even a hybrid individual between the two (*A. chinensis* and *A.*  
7 *macrotera*) as the other progenitor.

8 The AFLP study fails to resolve this issue because it does not indicate any admixture in the  
9 *Abelia x grandiflora* (Fig. 5); this suggests that *A. uniflora*, which was not sampled in the  
10 AFLP study, is probably the paternal parent. A lack of signal indicating the hybrid origin of  
11 *A. x grandifolia*, in the Structure analysis, is intriguing given the phylogenetic evidence but  
12 the involvement of *A. uniflora* will need to be determined conclusively in subsequent genetic  
13 studies.

14 A clear disjunction between *A. uniflora* and *A. macrotera* can be observed (Fig. 8), which is  
15 mainly due to its habitat requirements such as higher elevation for *A. macrotera* but also  
16 because *A. chinensis* forms an introgression barrier between *A. macrotera* in the west and the  
17 relictual zone of *A. uniflora* in the east. This is corroborated by its morphology which is  
18 similar to *A. x grandiflora* (Figs. 7 & 10):

- 19 • Variable number of sepals 2-4.
- 20 • Infundibuliform-bilabiate white corolla, with faint markings on lower lip.
- 21 • Slightly exerted stamens.

22 This study highlights the genetic distinction between *A. uniflora* and *A. macrotera*, two taxa  
23 that are often difficult to identify based on their morphology alone (Table 3, Figs. 6 & 7). The  
24 identity and origin of *A. x grandiflora*, a hybrid between *A. chinensis* and *A. uniflora* is

1 weakly supported and *Abelia uniflora* shows evidence of allopatric speciation due to  
2 hybridisation and introgression between *A. macrotera* and *A. chinensis*.

3

#### 4 ***Abelia schumannii*, hybridization in sympatric zones**

5 *Abelia schumannii* was introduced in cultivation after *A. uniflora* (Edinburgh  
6 collection Wilson E.H. 1230 collected in July 1908 in Sichuan, Wenchuan Xian). *Abelia*  
7 *schumannii* is now the only commonly available species of *Abelia* in the horticultural trade,  
8 and has been used in most selection programs.

9 *Abelia* ‘Edward Goucher’ is the hybrid *A. x grandiflora* x *schumannii*, obtained at Glenn  
10 Dale Plants, USA, in 1911. This is one of the oldest cultivars created and it is still commonly  
11 grown in the USA but it is much less frequent in the rest of the world. *Abelia* ‘Edward  
12 Goucher’ sepals are two and one is often notched at the apex, the corolla is similar to *A.*  
13 *schumannii* but the style is often exerted and the leaves are glossy adaxially. In its  
14 morphology it is similar to *A. uniflora* except the dark purple-pink corolla and glossy leaves  
15 (Figs. 9 & 10).

16 *Abelia* ‘Edward Goucher’ is identified as a hybrid in the AFLP study with admixture from *A.*  
17 *schumannii* and *A. x grandiflora* (Figs. 3--5).

18 *Abelia* ‘Maurice Foster’ was collected by Maurice Foster in Sichuan probably around Luding  
19 and was grown by Liss Forest Nurseries (UK). *Abelia* ‘Maurice Foster’ leaves are leathery,  
20 glossy adaxially and with prominent teeth on the margin and an acuminate apex. Flowers  
21 appear in short cymes or in clusters, with a purplish-pink corolla that has an inflated mouth. It  
22 superficially resembles *A. macrotera* var. *macrotera* but the leaves are smaller and the corolla  
23 mouth is more inflated (Fig. 6). The area where it was collected is covered by both *A.*

1 *macrotera* var. *zabelioides* and *A. schumannii* and many herbarium specimens from the same  
2 area have intermediate characters between the two taxa.

3 *Abelia* ‘Maurice Foster’ admixture was identified in STRUCTURE between *A. macrotera*  
4 and *A. schumannii* (Fig. 5). We can infer that hybridization, in sympatric regions of Sichuan,  
5 is common between taxa.

6 The AFLP study revealed that *Abelia schumannii* is a distinct species (Figs. 3 & 4) which  
7 was unexpected because the morphology is similar to *A. macrotera*. Its wide tube abruptly  
8 flaring to the corolla mouth is one of the most striking characters and probably linked to a  
9 specialized pollination syndrome and its distribution in the deep valleys of the QTP.

10 Backcrosses between closely related taxa such as *A. macrotera* and *A. schumannii* creates a  
11 similar effect to introgression but have a different morphological signature (Figs. 6 & 7):

- 12 • Leaves adaxially glossy.
- 13 • Number of sepals 2 with one notched apically.
- 14 • Bilabiate, inflated and purplish pink corolla with orange markings on lower lip.
- 15 • Inserted stamens, slightly exserted style.

16

### 17 ***Abelia forrestii* limited introgression and gene infiltration**

18 *Abelia forrestii* is the most localised taxon of *Abelia* only occurring in two localities  
19 within two of the deepest valleys of China. With its five sepals, long tubular-infundibuliform  
20 corolla, white-pink corolla without markings and fragrant flower it is one of the most distinct  
21 taxa in the genus (Figs. 6 & 7).

22 *Abelia* ‘Saxon Gold’ is also one of the more unusual cultivars obtained with its 5 sepals,  
23 bilabiate, purplish pink corolla without markings and yellow leaves (Figs. 9 & 10).

1 In the STRUCTURE analysis *A. 'Saxon Gold'* shows an admixture between *A. x grandiflora*  
2 and *A. schumannii*, indicating a backcrossed origin (Fig. 5). In the same analysis one  
3 individual of *A. forrestii* shows a small admixture with *A. macrotera*. It seems that the two  
4 taxa have a similar origin and that the isolation of *A. forrestii* within the deep valleys of the  
5 Nu-Jiang (Salween) and Yangtze Rivers allowed for limited introgression followed by  
6 allopatric speciation. Their morphological signatures are similar:

- 7 • Number of sepals 5.
- 8 • Infundibuliform pinkish white and scented corolla without markings on lower lip.
- 9 • Slightly exerted stamens and style.

10

11 ***Abelia macrotera* var. *deutziaefolia* and *A. chinensis* var. *lipoensis* chloroplast capture**

12 *Abelia macrotera* var. *deutziaefolia* is distributed in southern Yunnan, Guangxi and  
13 Guizhou provinces (Fig. 8). It has stems which are densely pubescent, its purple corollas have  
14 faint markings and have a short tube, the style is slightly exerted and the sepals are obtuse  
15 (Figs. 6 & 7).

16 *Abelia* 'II6306' is an experimental hybrid sent by Liss forest Nursery (UK) it resembles *A.*  
17 *chinensis* but its inflorescence are loose and corollas are bilabiate-infundibulform as in *A. x*  
18 *grandiflora* (Fig. 10).

19 In the STRUCTURE analysis *A. 'II6306'* shows admixture between *A. x grandiflora* and *A.*  
20 *chinensis*, indicating a backcrossed origin (Fig. 5).

21 Organellar DNAs (i.e., chloroplast DNA and mitochondrial DNA) can be used for tracing the  
22 long-term effects of hybridization in natural populations. Because their inheritance is  
23 uniparental, groups of associated loci are not separated by recombination, so that a great deal  
24 of historical information is preserved in these sequences (Whittemore and Schaal, 1991). As

1 seen in the Bayesian tree of *Abelia* based on the combined *rbcL*, *trnL-F* and *matK* sequence  
2 data (Fig. 1), *Abelia* x *grandiflora*, *Abelia chinensis* and *A. macrotera* var. *deutziaefolia* form  
3 a polytomy (BP 54/ PP 0.96). Hybridization followed by introgression between *A. chinensis*  
4 and *A. macrotera* can therefore be hypothesised.  
5 *Abelia macrotera* var. *deutziaefolia* and *Abelia chinensis* var. *lipoensis* both share a similar  
6 distribution range in southern Yunnan, Guizhou and Guangxi where both species *Abelia*  
7 *macrotera* and *Abelia chinensis* grow sympatrically. *Abelia* 'II6306' and *Abelia chinensis*  
8 var. *lipoensis* show a similar loose inflorescence as well as other morphological characters  
9 associated with *A. macrotera* (Fig. 7):

- 10 • Number of sepals 5.
- 11 • Loose inflorescence.
- 12 • Glabrous stems.
- 13 • Infundibuliform-bilabiate and white corollas.
- 14 • Stamens exserted.

15 *Abelia macrotera* var. *deutziaefolia* is on the other end of the spectrum, being introgressed  
16 from *A. chinensis* and sharing a few similar morphological characters (Figs .6 & 7).

- 17 • Number of sepals 2.
- 18 • Bilabiate purplish pink corolla with faint orange markings on lower lip.
- 19 • Inserted stamens, slightly exserted style.

20

#### 21 ***Abelia chinensis* var. *hanceana* segregation across Japan vs. eastern China**

22 *Abelia chinensis* var. *hanceana* is distributed in Xiamen and Taiwan, China as well as  
23 the Ryukyu Islands but not reaching Kyushu, Japan. This variety is characterised by its

1 smaller size and compact inflorescences. Its leaves have a crenate margin and resemble  
2 leaves of *Abelia uniflora* (Figs. 6 & 7).  
3 *Abelia* 'Rose Creek' is a backcross *A. x grandiflora* x *A. chinensis* and resembles *Abelia*  
4 *chinensis* but it has more compact inflorescences (Figs. 9 & 10). In the STRUCTURE analysis  
5 *A. 'Rose Creek'* also showed admixture between *A. x grandiflora* and *A. chinensis* indicating  
6 a backcrossed origin (Fig. 5). The Okinawa trough and associated straits began to rift at  
7 1.55 Ma, isolating the Ryukyu Islands from the Chinese continent, Japanese islands, Taiwan  
8 Island and some of the islands from each other. Physical isolation began to generate the  
9 allopatric speciation within these islands (Osozawa *et al.*, 2015). Vicariance of *Abelia*  
10 *chinensis* var. *hanceana* across the East China Sea between Xiamen, Taiwan and the Ryukyu  
11 Islands may have favoured limited backcrossing along the eastern edge of *Abelia chinensis*  
12 range.

13 Some characters shared are (Fig. 7):

- 14 • Number of sepals 5.
- 15 • Compact inflorescence.
- 16 • Infundibuliform, white corollas.
- 17 • Stamens long exserted.

18

### 19 ***Hybridization-backcrossing driven speciation***

20 We have highlighted the diversity of the genus and clarified the status of some of the  
21 most controversial names. The confusion between names is probably due to the complex  
22 evolutionary histories between taxa, the principal forces being hybridization and  
23 introgression. Breeding in horticulture has merely reproduced natural processes that occur in  
24 longer time scales and has been blurred by introgression. Character infiltration due to back  
25 crossing is well known to horticulturists and is one of their major aims, this phenomenon also

1 provides an evolutionary advantage that leads to novel adaptations and cryptic taxa. Some of  
2 the infiltrated characters are here identified (Table 6, Fig. 11):

- 3 • Calyx lobe numbers 2, 2-5 or 5
- 4 • Inflorescence compact or loose
- 5 • Leaves glossy adaxially.
- 6 • Stems glabrous.
- 7 • Corolla bilabiate, infundibulform or bilabiate-infundibulform or inflated.
- 8 • Flowers white, pinkish purple with or without orange markings.
- 9 • Flower scented.
- 10 • Exserted stamens and style.

11 *Abelia* 'Edward Goucher' has a similar origin to *A.* 'Saxon Gold' but exhibits very different  
12 characters only having two sepals a bilabiate purplish-pink corolla with strong markings and  
13 an exserted style. This demonstrates that recombination during hybridisation is random and  
14 can produce new combinations of genes which in turn can increase the morphological  
15 diversity and result in novel characters belonging to different species. Sepal number, corolla  
16 shape and inflorescence structure are amongst these easily identifiable characters in *Abelia*  
17 that could be transferred between taxa. The establishment of new intraspecific taxa is a  
18 frequent outcome of introgression; backcrossing to one or both parents leads to the  
19 infiltration of specific genes from one species to another (Fig. 11). Such interspecific gene-  
20 flow, known as introgression, results in the production of offspring that are clearly referable  
21 to one of the parent species, but that possess certain characters inherited from the second  
22 species. Sometimes these products may become stabilised and develop into a new  
23 intraspecific taxon or cryptic species ([Abbott, 1992](#)).

24 Several cryptic taxa are here identified (Table 6):



- *Abelia chinensis* backcrosses with: *A. chinensis* var. *hanceana* (possibly with *A. uniflora*) and var. *lipoensis* (possibly with *A. macrotera* var. *macrotera*) / *Abelia* ‘Rose Creek’, *Abelia* ‘II6306’.
- *Abelia uniflora* s.l. backcrosses with: *A. macrotera* var. *deutziaefolia* and *A. uniflora*, *A. forrestii* / *Abelia* ‘Edward Goucher’, *Abelia* ‘Saxon Gold’.

Backcrossing does not necessarily occur exclusively between two taxa and can involve many taxa belonging to the same complex such as the *A. uniflora* species complex and *A. chinensis*. This is less likely in the wild because it requires taxa to be growing sympatrically.

## CONCLUSIONS

‘It appears to be necessary to accept that some taxa are highly variable in the wild and that, consequently, some names in common use in the West merely represent points in a continuous spectrum of variation of what is better regarded as a single species’ Barnes, P (2001). Our aim was to bridge the gap between horticulturalists and the botanical community in order to understand the genetic diversity within *Abelia* and where to concentrate efforts into conserving and exploiting the benefits offered by ornamental plants. A clearer view of the taxonomy, speciation mechanisms and morphology is here presented and highlights the importance of hybridization and backcrossing in breeding programs and also in natural speciation. It is not surprising that with these types of complex and multidirectional processes, taxa of *Abelia* are often difficult to identify and the species boundaries are blurred. The knowledge of the genetic diversity and genetic relatedness within *Abelia* is potentially useful to improve the current strategies in breeding and germplasm conservation to enhance the ornamental and economic value of the genus. An understanding of both the genetic diversity and the population structure of *Abelia* in China can also provide insight into the conservation and management of some endangered taxa.

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1    **Table 1.** Statistics of the three regions used

	ITS	<i>rbcL</i>	<i>trnL-F</i>	<i>matK</i>	CpDNA Combined
No. taxa	18	24	24	24	24
Total aligned characters	637	604	849	779	2232
Constant characters	572	541	736	634	1911
Parsimony informative	40	61	52	76	153
Consistency index	0.814	0.57	0.923	0.889	0.833
Retention index	0.91	0.484	0.945	0.923	0.865
Evolution model	GTR+G	HKY+G	GTR+G	GTR+G	-

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3    **Table 2.** Statistics of the AFLP analysis

	G13 nr.	G13 %	G2 nr.	G2 %	Y15 nr.	Y15 %
Initial bin number	149.00	100	166.00	100.00	173.00	100.00
Final bin number	76.00	51.01	111.00	66.87	113.00	65.32
Removed low intensity	49.00	32.89	44.00	26.51	50.00	28.90
Removed non replicable bin	21.00	14.09	8.00	4.82	10.00	5.78
Removed rare Frequency bin	3.00	2.01	3.00	1.81	0.00	0.00
Error rate bin	0.0658	0.00	0.0360	0.00	0.0487	0.00
Ibin	0.2824	0.00	0.2666	0.00	0.2485	0.00

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1     **Table 3.** Morphological characters of the five species of *Abelia*

Characters	<i>Abelia chinensis</i>	<i>Abelia schumannii</i>	<i>Abelia uniflora</i>	<i>Abelia macrotera</i>	<i>Abelia forrestii</i>
<b>Corolla shape</b>	infundibuliform-campanulate	bilabiate with inflated mouth	bilabiate with inflated mouth	bilabiate	tubular-infundibuliform
<b>Corolla colour</b>	white without markings	purplish-pink with orange markings	white-Purplish with faint yellow markings	purplish-pink with orange markings	purplish-pink without markings
<b>Fragrance</b>	scented	Non-scented	?	Non-scented	scented
<b>Bloom</b>	Sept--Dec	May--Sept	May--Sept?	May--Sept	May--Sept
<b>Anthers and style</b>	long exserted	inserted	inserted to exserted	inserted to slightly exserted	inserted to slightly exserted
<b>Calyx lobes</b>	5	2	2--4	2	5
<b>Flowers</b>	paired	single	single	single	single
<b>Epicalyx bracts</b>	6	4	4	4	4
<b>Inflorescence</b>	terminal, flowers many and cymose	flowers few axillary	flowers few axillary	flowers few axillary	flowers few axillary

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3     **Table 4.** Morphological characters of the cultivated species of *Abelia*

Key characters	<i>Abelia chinensis</i>	<i>Abelia x grandiflora</i>	<i>Abelia uniflora</i>	<i>Abelia schumannii</i>
Leaves	glossy adaxially	glossy adaxially	?	not glossy
Leaves	semi-evergreen	semi-evergreen	?	deciduous
Corolla shape	infundibuliform-campanulate	infundibulform-bilabiate	bilabiate	bilabiate
Corolla colour	white without markings	purplish-pink with orange markings	white-pink with faint yellow markings	purplish-pink with orange markings
Fragrance	scented	scented	?	non scented
Bloom	autumn	summer-autumn	summer	summer
Androecium	long exserted	inserted to exserted	inserted	inserted
Style	long exserted	slightly exserted	slightly exserted	inserted
Calyx	5	2--5	2--4	2
Flowers	paired opening consecutively	single	single	single
Epicalyx	6	4	4	4
Inflorescence	terminal compact	loose to compact	loose	loose

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8     **Table 5.** List of *Abelia* Cutivars names and descriptions.

Cultivars name	Preferred selling name/ synonyms	Breeder	Date	Parentage	Description
‘Abghop’	HOPLEYS	Hopleys Nurs.	1992	'Francis Mason' Sport	Dwarf, variegated leaves, teratological white flowers.
‘Bumble Bee’		?	?	<i>Abelia schumanii</i>	Selection with larger flowers.
‘Canyon Creek’		Dirr, M.	2002	<i>A. chinensis</i> x <i>A. x grandiflora</i>	Sepals 4--5, corolla infundibuliform-bilabiate pink without markings, style and stamens not exerted inflorescence not dense terminal leaves green, glossy.
‘Compact’	‘Compacta’	?	?	similar to 'Sherwood'	The white flowers have a basal spur at the base like the rest of the 'Sherwood' group.
‘Conti’	CONFETTI	Flowerwood Nurs.	1987	Sport of 'Sherwood'	Variegated dwarf form with teratological flowers.
‘Edward Goucher’		Glenn Dale Plants	1911	<i>A. x grandiflora</i> x <i>schumanii</i>	Flowers single, bilabiate, axillary dark purple. Sepals 2 often notched at the apex. Leaves glossy.
‘Francis Mason’		Mason nursery New Zealand	1950 or 1970	<i>A. x grandiflora</i> selection	Variegated form of <i>A. x grandiflora</i> . Copper coloured young shoots. Leaves variegated yellow or yellow margined.
‘Goldenglossy’	GOLD DUST	Jerry Brunson.		Sport of 'Sherwood'	Compact with yellow variegated leaves.
‘Goldspot’ group				Selections of <i>A. x grandiflora</i> 'Francis Mason'	With more stable and less reversion or more yellow variegation.
‘Kaleidoscope’					
‘Kaleidoscope’		Plant Haven Inc.	1997	Sport of A. ‘Little Richard’	Dwarf, leaves with cream yellow and orange variegation.
‘Keiser’	RUBY ANNIVERSARY	Susan Keiser	1999	Back cross chinensis	Not so compact inflorescences. Sepals 4--5, corolla infundibuliform white without markings, style and stamens exerted leaves green, glossy.
‘Lavender Mist’		University of Georgia	2006	<i>A. 'Edward Goucher' x Abelia chinensis</i>	Heavy bloomer, with clusters of fragrant lavender flowers. Sepals 5 reddish. (HortScience 41:967-1084 (2006))
‘Little Richard’		Curran's nursery	?	Sport of 'Sherwood'	Lustrous dark green leaves. Large flowers, corolla whitish with 1--2 spurs at the base of the corolla, typical of Sherwood group.
‘Lynn’	PINKY BELLS	Spring meadow Nursery	2003	<i>Abelia</i> 'Bumblebee' x <i>A. 'Little Richard'</i>	Compact plant with large pendulous, lavender-pink flowers. Sepals mostly 2, corolla almost infundibuliform pink with large broad mouth, markings faint.

‘Mardi Gras’		Crowder Rick	2003	<i>Abelia x grandiflora</i> x <i>chinensis</i> sport	Leaves green with white and pink variegated margins.
‘Maurice Foster’		Liss Nursery ex. 19770008	1977	<i>Abelia macrotera</i> x <i>A. schumanii</i>	Collected by Maurice Foster in Sichuan (probably Luding). Inflorescence in short axillary raceme, sepals narrow with acute apex, Leaves leathery.
‘Mei-fu-hana-tsukubane-utsugi’		Suda, H. 1976. Variegated plants :180	1972	Irradiated <i>A. x grandiflora</i>	Yellow margined and blotched leaves. Obtained by gamma-rays.
‘Minduo1’	SUNNY ANNIVERSARY	Olivier Nazeyrollas, Beaufort en Vallee.	2005	<i>A. x grandiflora</i> ‘Minfest’ x unknown	Sepals mostly 2, corolla almost infundibuliform pink with large broad mouth, markings strong reaching base of corolla tube. Style exserted.
‘Panache’ or ‘Panash’	‘Silver Panache’			<i>A x grandiflora</i> ‘Prostrata’	Stable variegated leaves, silver to yellow margined. Prostate, Flowers teratological like ‘Conti’.
‘Pleasant Surprise’		Pleasant View Nurs.	2002	<i>A. x grandiflora</i> x <i>chinensis</i> .	(Hort Science 38(6): 1300. 2003). Flowers single, sepals 5, corolla infundibuliform-bilabiate, pinkish white, sepals obovate, Inflorescence dense.
‘Plum Surprise’		University of Georgia	2006	<i>A. 'Edward Goucher'</i> x <i>A. 'Francis Mason'</i>	(HortScience 41:967-1084 (2006) Sepals 4-5, Corolla bilabiate white-pink with faint markings. Leaves small moderately glossy and green.
‘Prostrate White’	‘Prostrata’	Abbott, D.	1955	Similar to <i>A. ‘Sherwood’</i>	Similar to <i>A. ‘Sherwood’</i> but more prostrate.
‘Radiance’		Plant Haven Nurs.	2006	<i>A. ‘kaleidoscope’</i>	Variegated compact sport.
‘Raspberry Profusion’		University of Georgia	2006	<i>A. 'Edward Goucher'</i> x <i>Abelia chinensis</i>	(HortScience 41:967-1084 (2006)). Very heavy bloomer. Panicles are large and showy with fragrant pink flowers and raspberry-colored sepals.
‘Rika 1’	BRONZE ANNIVERSARY	Grand Haven	2003	<i>Abelia x grandilora</i> ‘Minipan’	Young leaves with bronze colour, similar to ‘Francis Mason’.
‘Rose Creek’		Dirr, M.	2001	<i>A. chinensis</i> x <i>A. x grandiflora</i>	Sepals 5, corolla infundibuliform white, style and stamens exserted inflorescence very dense terminal leaves glossy.
‘Saxon Gold’		Saxon Nurs.	1997	<i>Abelia schumanii</i> <i>A. x grandiflora</i> ?	This is a very intriguing cultivars, flowers and inflorescence are like <i>Abelia schumanii</i> but differs in having 5 sepals and golden yellow foliage.

<b>‘Sherwood’ or ‘Sherwoodii’</b>	Published as ‘Sherwoodi’.		1935	Sport of <i>A. x grandiflora</i>	American Assoc. Nurs. Proc. 1949:123. Very compact.
<b>‘Short and Sweet’</b>		Robert Pearce	2000	<i>A. x grandiflora</i> ‘Compacta’	(Plant Variety journal 14(1):15.2001). similar to <i>A. ‘Conti’</i> .
<b>‘Sunrise’</b>		Taylor’s nursery NC, USA	1996	<i>A x grandiflora</i> mutation	Smaller but not as ‘Conti’. Leaves yellow margined to cream orange or red margined, stem dark red. Teratological flowers.
<b>‘White Marvel’</b>		Sakaue; Katsuya, Hines Nurs.	2000	<i>A. x grandiflora</i> ‘Francis Mason’	New shoots and foliage which emerge white, cream, pale yellow to gold with occasional terminal golden orange foliage.

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**Table 6.** Morphological character changes paralleled between cultivars and wild taxa.

	← + introgression -			- introgression + →	
	Backcross ' <i>uniflora</i> '	Backcross ' <i>uniflora</i> '	Hybrid	Backcross <i>chinensis</i>	Backcross <i>chinensis</i>
<b>Sepals</b>	2	2--3	2--5	4--5	5
<b>Epicalyx</b>	4	4	4	4 or 6	6
<b>Flower</b>	single	single	single	single-paired	paired
<b>Inflorescence</b>	loose	loose to compact	loose to compact	loose	dense
<b>Corolla</b>	bilabiate	infundibuliform-bilabiate or bilabiate	infundibuliform-bilabiate	infundibuliform-bilabiate or infundibuliform-campanulate	infundibuliform-campanulate
<b>Fragrance</b>	non-scented	scented or non-scented	scented	scented	scented
<b>Bloom</b>	summer	summer-autumn	summer-autumn	summer-autumn	autumn
<b>Androecium and style</b>	inserted	inserted to exerted	inserted to exerted	inserted to exerted	long exerted
<b>Wild taxa</b>	<i>A.uniflora</i>	<i>A.forrestii</i>		<i>A. chinensis</i> var. <i>lipoensis</i>	<i>A.chinensis</i> var. <i>hanceana</i>
<b>Cultivated taxa</b>	A.'Edward Goucher'	A.'Saxon Gold'	A.x grandiflora	A.'II6306'	A.'Rose Creek'

**Figure 1.** Bayesian tree of *Abelia* and outgroups based on the combined *rbcL*, *trnL-F* and *matK* sequence data. MP (first), bootstrap branch support and Bayesian posterior probabilities (last) indicated above a cut-off value of 50 and 0.5, respectively.

**Figure 2.** Bayesian tree of *Abelia* and outgroups based on the Internal Transcribed Spacer (ITS) sequence data. MP (first), bootstrap branch support and Bayesian posterior probabilities (last) indicated above a cut-off value of 50 and 0.5, respectively.

**Figure 3.** Neighbour Joining Tree of 29 *Abelia* cultivars and taxa based on AFLP analysis with three pair primer combinations

1 **Figure 4.** Principal coordinates analysis based on the distance between individual amplified  
2 fragment length polymorphism (AFLP)

3 **Figure 5.** STRUCTURE analysis of *Abelia* cultivars and taxa (K=2, 4 and 5)

4 **Figure 6:** Photographs of *Abelia* taxa. A--B. *Abelia macrotera* var. *deutziaefolia* from  
5 Guizhou, Guiyang note the short thick tube and exserted style; C. *Abelia uniflora* cultivated  
6 in Wuhan B.G. note the white corolla and faint yellow markings; D & G. *Abelia chinensis*  
7 var. *chinensis* cultivated at Kew, note the campanulate corolla and long exserted styles and  
8 stamens; E. *Abelia forrestii* var. *forrestii* from Yunnan, Nujiang, note the long tubular,  
9 infundibuliform corolla without markings; F. *Abelia macrotera* var. *mairei* from Yunnan,  
10 Kunming; H. *Abelia schumanii* cultivated, note the large and wide corolla mouth; I. *Abelia*  
11 *macrotera* x *A. schumanii* ‘Maurice Foster’ cultivated; J. *Abelia macrotera* var. *macrotera*  
12 from Sichuan, Bazhong.

13 **Figure 7:** Line drawings illustrating key morphological characters of *Abelia* taxa. A. Flowers  
14 in lateral views; B--G. *Abelia forrestii* var. *forrestii*; B. corolla; C. Nectary bulge; D. Corolla  
15 mouth; E. Longitudinal section of the corolla; F. Androecium; G. Style; H. Flowering  
16 branches; I. Epicalyx and achenes; J. Habit; K--M & O--P. *Abelia macrotera* var.  
17 *deutziaefolia*; K. Stem node indumentum; L. Adaxial leaf, M. Abaxial leaf; N. *Abelia*  
18 *chinensis* var. *hanceana* adaxial leaf; O. Abaxial leaf surface; P. Adaxial leaf surface.

19 **Figure 8:** Distribution map of *Abelia* taxa drawn from herbarium specimens using BRAHMS  
20 and DIVA GIS. Map obtained from 761 collections.

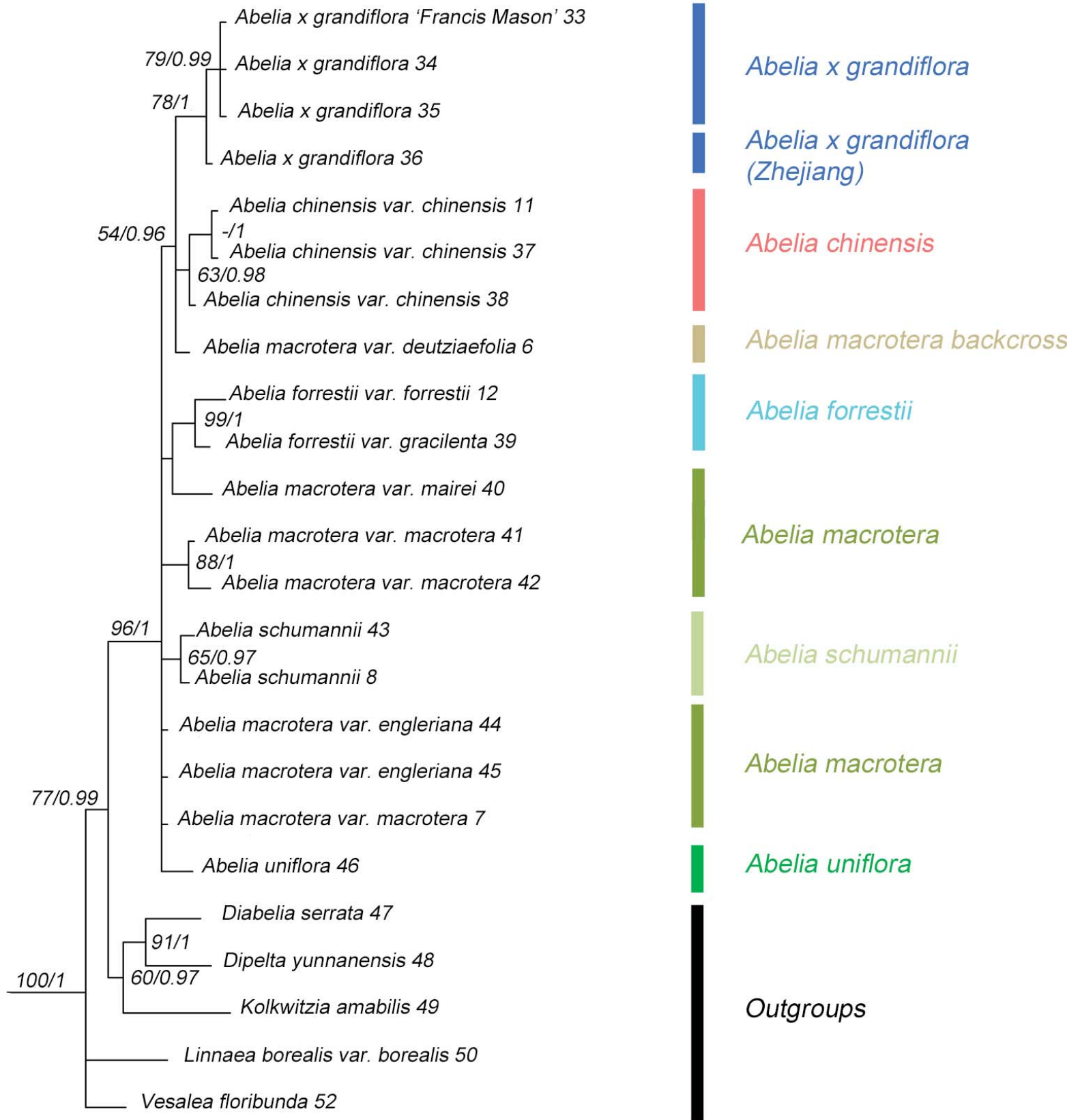
21 **Figure 9:** Photographs of *Abelia* cultivars at the RHS Wisley trial garden A--B. *Abelia*  
22 ‘Edward Goucher’ note the two sepals and the notched apex; C--D. *Abelia* ‘Saxon Gold’ note  
23 the five sepals and yellow leaves; E--F. *Abelia* x *grandiflora* ‘Francis Mason’; G. *Abelia* x

1 *grandiflora* ‘Conti’ note the variegated leaves; H. *Abelia x grandiflora* ‘Sherwood’ note the  
2 basal spur on the corolla; I--J. *Abelia* ‘Pleasant Surprise’; K. *Abelia* ‘Rose Creek’ note the  
3 dense terminal inflorescence and campanulate corolla.

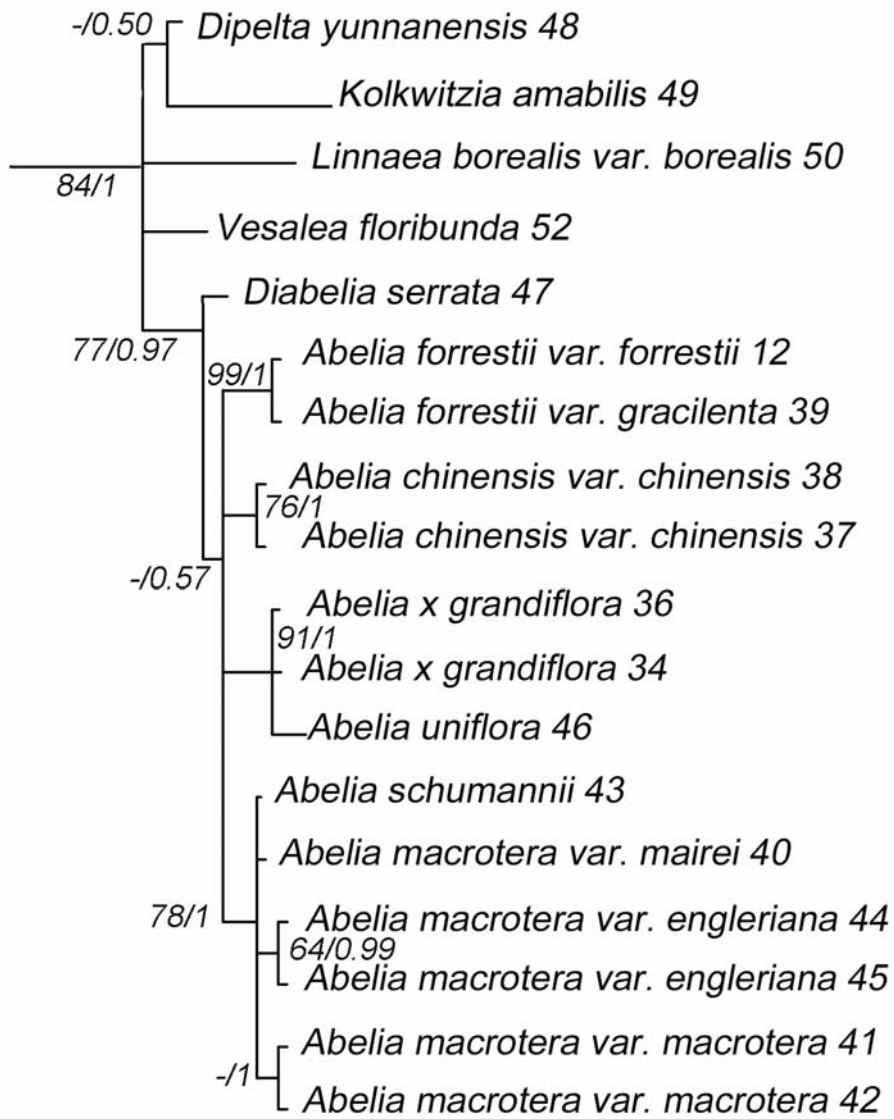
4 **Figure 10:** Line drawings illustrating key morphological characters of *Abelia* cultivars. A.  
5 Flowers in lateral views; B--G. *Abelia x grandiflora* ‘Francis Mason’; B. corolla; C. Nectary  
6 bulge; D. Corolla mouth; E. Longitudinal section of the corolla; F. Style; G. Androecium; H.  
7 Flowering branches; I. Epicalyx and achenes; J. Habit; K--O. *Abelia x grandiflora* ‘Francis  
8 Mason’; K. Stem node indumentum; L. Adaxial leaf, M. Abaxial leaf; N. Abaxial leaf  
9 surface; O. Adaxial leaf surface.

10 **Figure 11:** Principal coordinates analysis based on the distance between individual amplified  
11 fragment length polymorphisms (AFLPs) plotted with two of the main diagnostic characters  
12 for the genus: sepal number and inflorescence with single or paired flowers.

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Outgroups

*Abelia forrestii*

*Abelia chinensis*

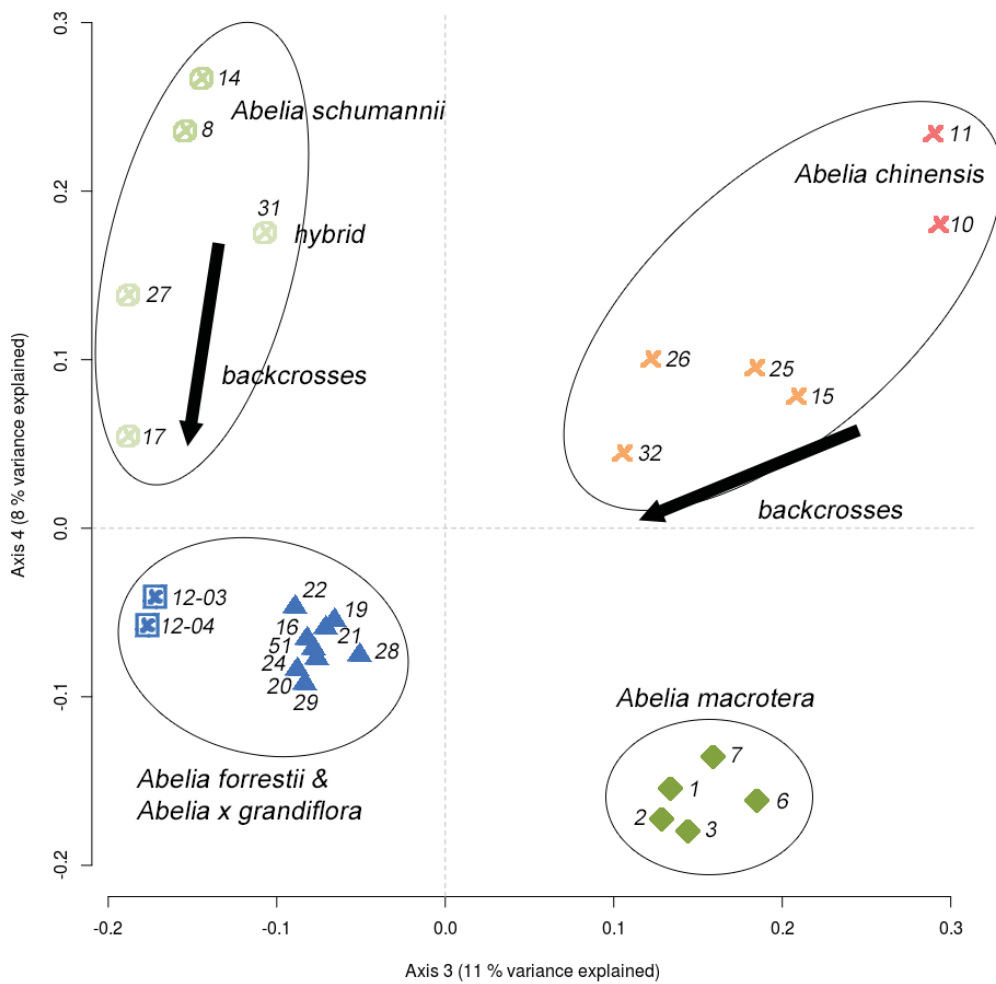
*Abelia x grandiflora*

*Abelia uniflora*

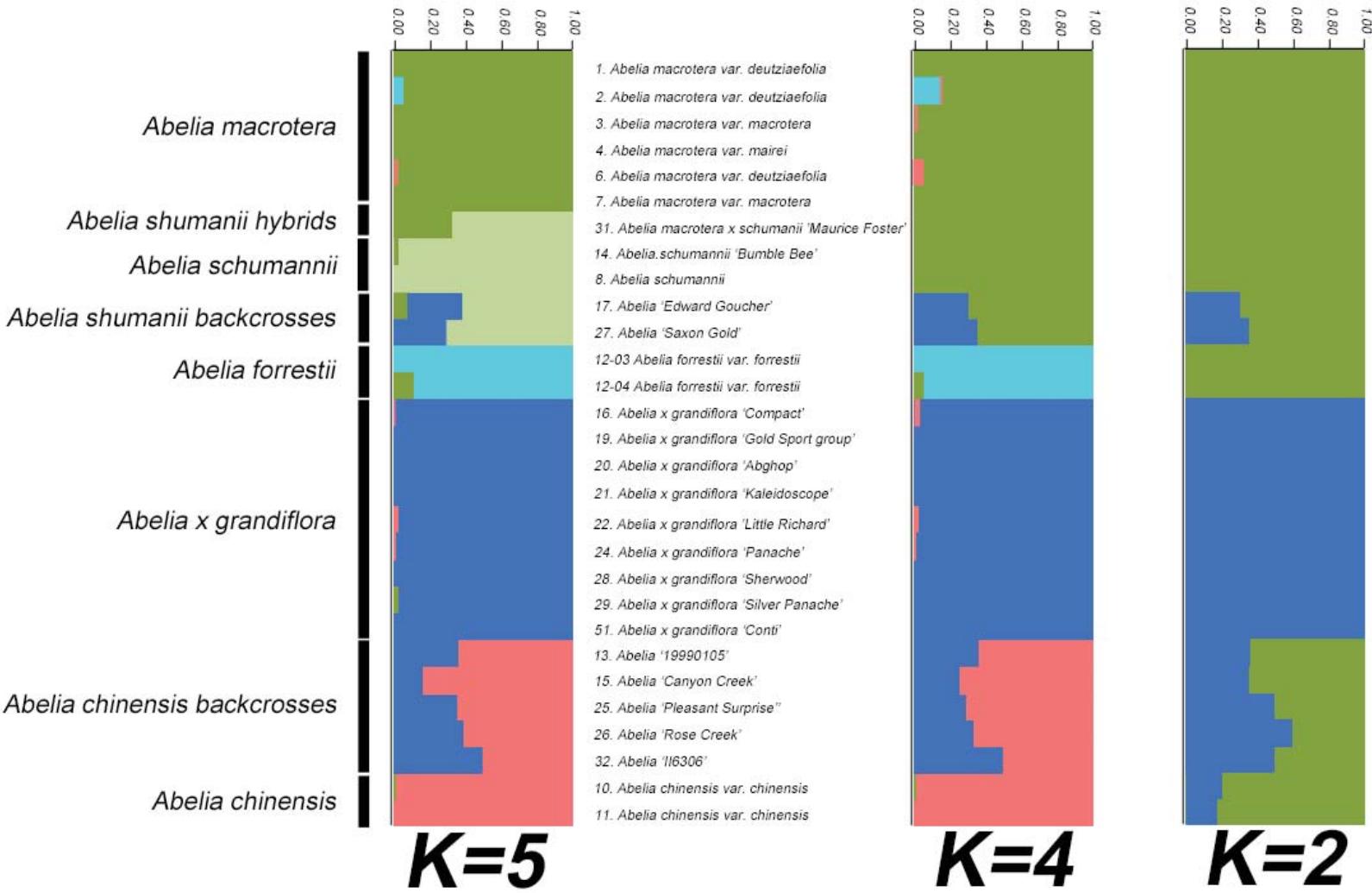
*Abelia schumannii*

*Abelia macrotera*





- ⊗ *Abelia schumannii*
- ⊗ *Abelia shumannii* hybrids and backcrosses
- × *Abelia chinensis*
- × *Abelia chinensis* backcrosses
- ◆ *Abelia macrotera*
- ▲ *Abelia x grandiflora*
- ⊠ *Abelia forrestii*







**A**



**B**

***Abelia macrotera*  
var. *deutziaefolia***



**C**

***Abelia uniflora***



**D**

***Abelia chinensis* var. *chinensis***



**E**

***Abelia forrestii***



**F**

***Abelia macrotera* var. *mairei***



**G**



**H**

***Abelia schumannii***



**I**

***Abelia* 'Maurice Foster'**



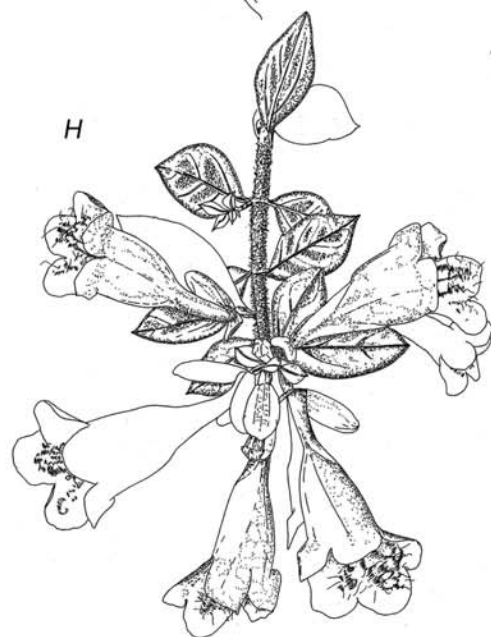
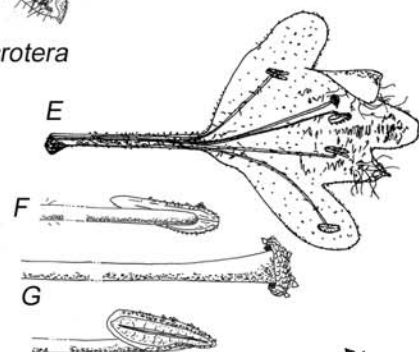
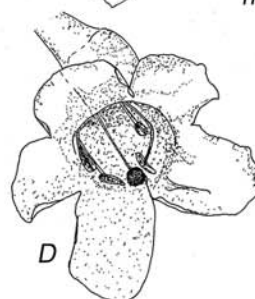
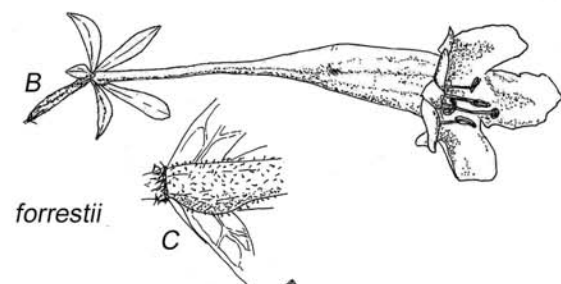
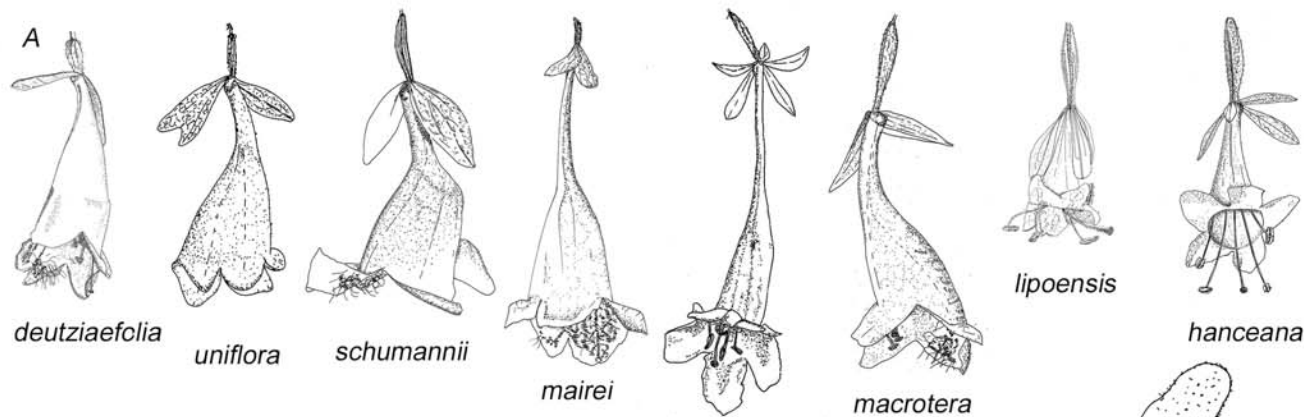
**J**

***Abelia macrotera* var. *macrotera***

*A. macrotera* back crosses

*A. forrestii*

*A. chinensis* back crosses



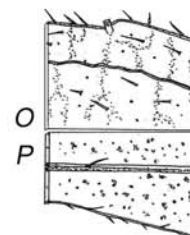
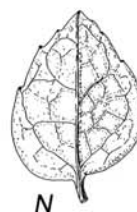
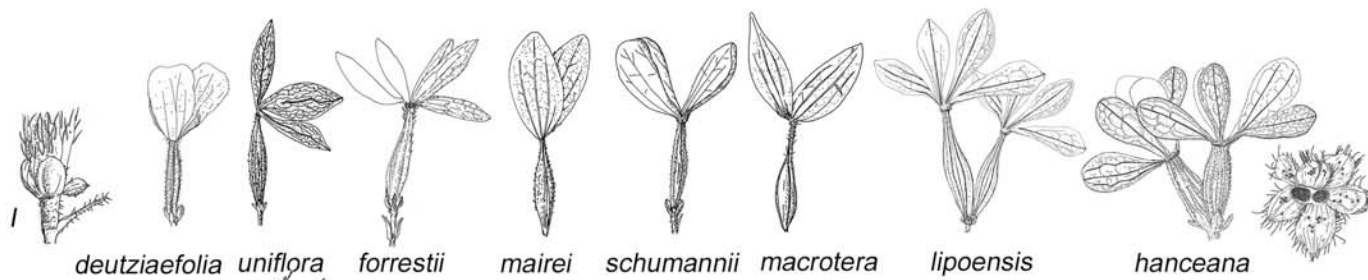
*deutziaefolia*



*forrestii*

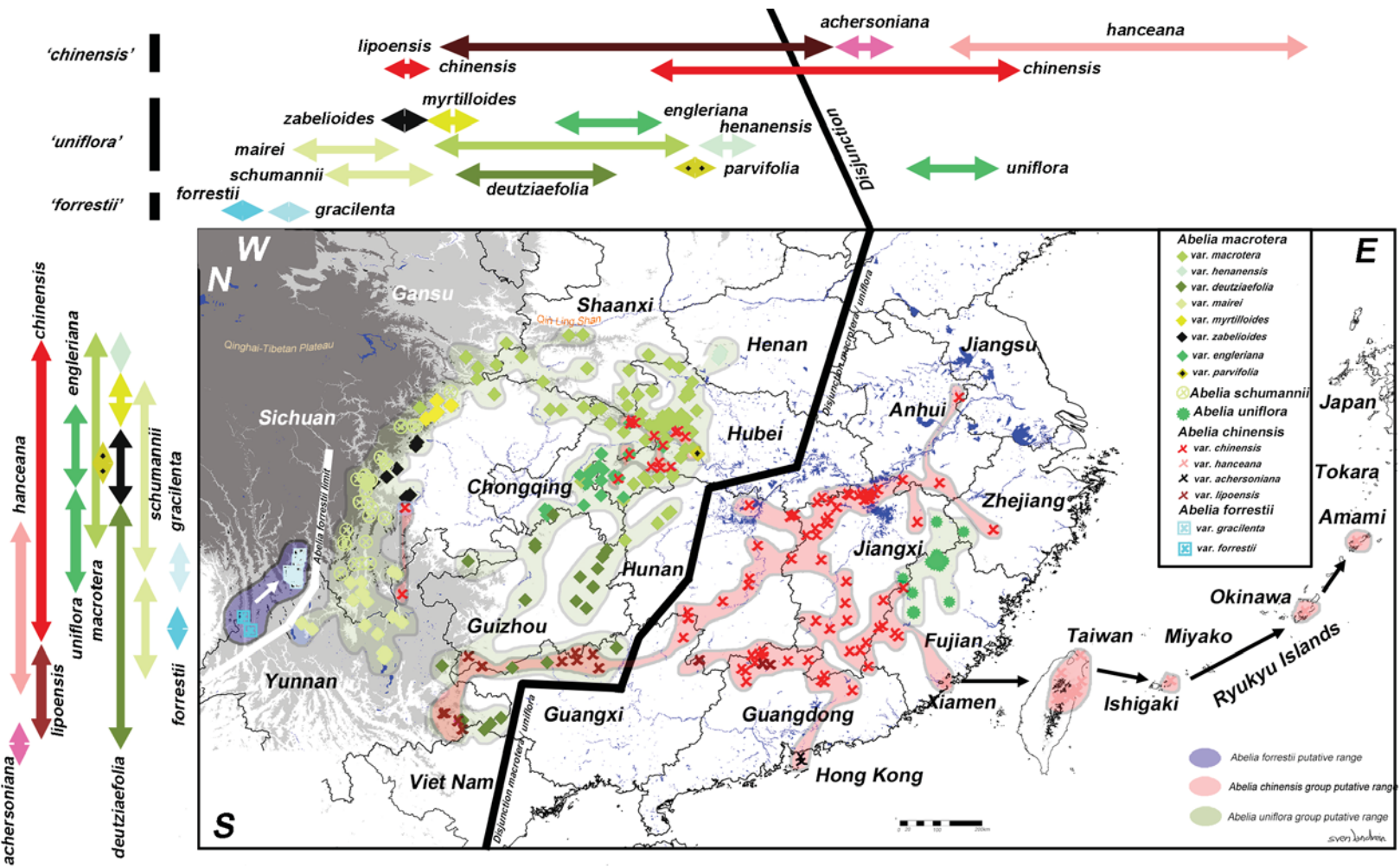


*hanceana*

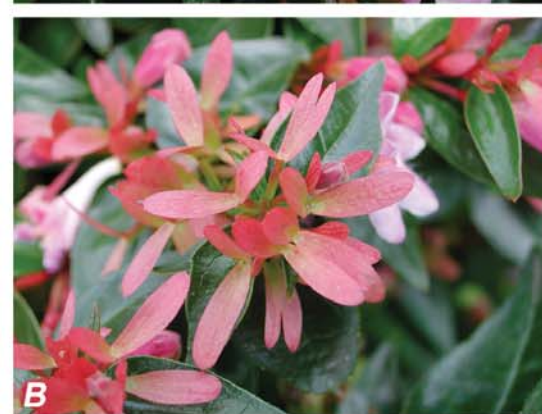
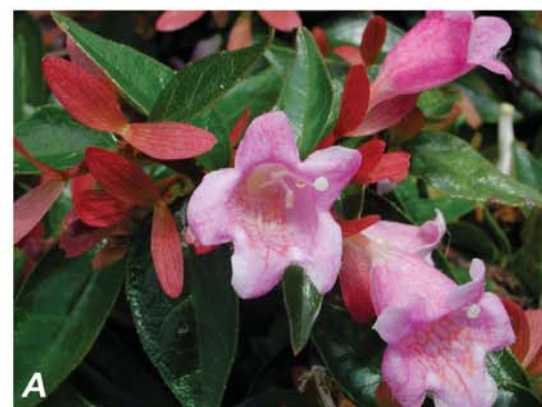


sven landren









*Abelia schumannii* backcrosses



*Abelia x grandiflora*



*Abelia chinensis* backcrosses



*A. schumannii* back crosses ← *A. x grandiflora* → *A. chinensis* back crosses

